Secreted IL-1ra

GAATTCCGGGCTGCAGTCACAGAATGGAAATCTGCAGAGGCCTCCGCAGTCACCTAATCACTCTCCTCTTCCTGTTCCATTCAG MetGluIleCysArgGlyLeuArgSerHisLeuIleThrLeuLeuPheLeuPheHisSer

Intracellular IL-1ra type I

IRA 1

CAGAAGACCTCCTGTCCTATGAGGCCCTCCCCATGGCTTTAG

Intracellular IL-1ra Type II

IRA 1

IRA 5

CAGAAGACCTCCTGTCCTATGAGGCCCTCCCCATGGCTTTAGCTGACTTGTATGAAGAAGGAGGTGGAGGAGGAGGAGGAGAAGGTGAAGACAATGCTGACTCAAAGG MetalaLeualaaspLeuTyrGluGluGlyGlyGlyGlyGlyGluGlyGluBspasnAlaaspSerLys

Common IL-1ra sequence

<u>AGACGATCTGCCGACCCTCTGGGAGAAAATCCAGCAAGATGCCAAGCCTTCAGAATCTGGGATGTTAACCAGAAGACCTTCTATCTGAGGACAACAACTAGTT</u> GluthrileCysArgProSerGlyArgLysSerSerLysMetGlnAlaPheArgIleTrpAspValAsnGlnLysThrPheTyrLeuArgAsnAsnGlnLeuVal AlaglyTyrLeuGlnGlyProAsnValAsnLeuGluGluLysIleAspValValProIleGluProHisAlaLeuPheLeuGlyIleHisGlyGlyLysMetCys

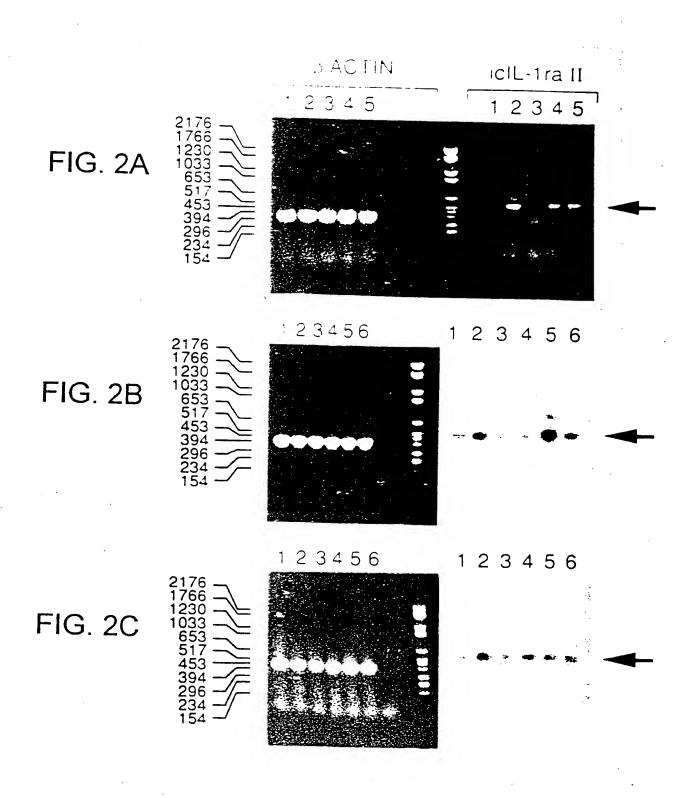
CTGTCCTGTGTCAAGTCTGGTGATGAGACCAGACTCCAGCTGGAGGCAGTTAACATCACTGACCTGAGCGAGAACAGAAAGCAGGACAAGCGCTTCGCCTTCATC LeuSerCysValLysSerGlyAspGluThrArgLeuGlnLeuGluAlaValAsnIleThrAspLeuSerGluAsnArgLysGlnAspLysArgPheAlaPheIle

ArgSerAspSerGlyProThrThrSerPheGluSerAlaAlaCysProGlyTrpPheLeuCysThrAlaMetGluAlaAspGlnProValSerLeuThrAsnMet

CCTGACGAAGGCGTCATGGTCACCAAATTCTACTTCCAGGAGGACGAGTAGTAC ProAspGluGlyValMetValThrLysPheTyrPheGlnGluAspGlu\*









1 2 3

— 97,4 — 66

<del>--- 45</del> ,

<del>---</del> 31

<del>---</del> 21,5

<del>-</del> 14,5



0.1 ng/m1 IL-1

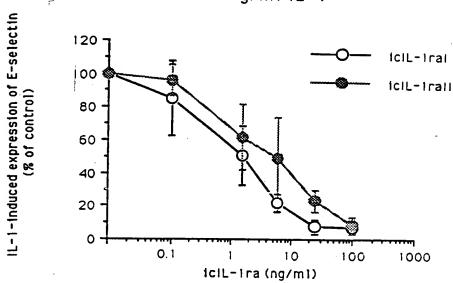
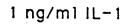


FIG. 4A



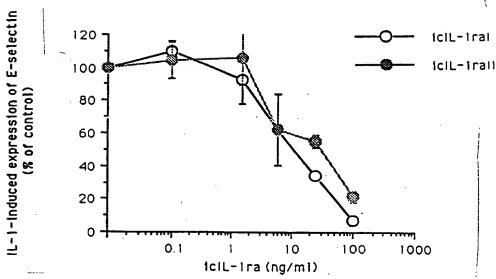


FIG. 4B

88-910,733

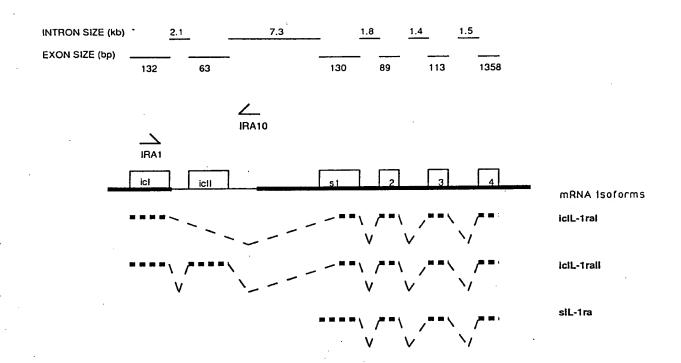
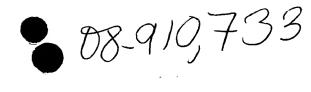


FIG. 5



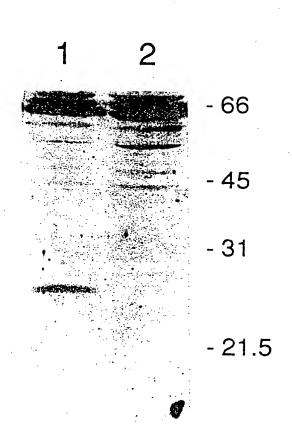


FIG. 6